

SEQUENCE LISTING

BARBOUR, ALAN G.

CARTER, CAROL <120> A DIAGNOSTIC TEST FOR INFECTION WITH A SPIROCHETE BORNE BY AMBLYOMMA AMERICANUM <130> UTSK:352USC1 <140> 10/620,795 <141> 2003-07-14 <150> 08/437,013 <151> 1995-05-08 <150> 09/275,506 <151> 1999-03-24 <160> 38 <170> PatentIn Ver. 2.1 <210> 1 <211> 641 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Primer <400> 1 acatattcag atgcagacag aggttctatt caaattgaaa ttgaacaact tacagatgaa 60 attaacagag ttgctgatca ggctcaatac aaccagatgc atatgttatc taacaaatca 120 tctgctcaaa atgtaaaaac tgctgaagag cttggaatgc aacctgcaaa aattaataca 180 ccagcatcac taactggagc acaagcttca tggacattga gagttcaggt aggtgcaaat 240 caggatgaag caattgctgt taatattttc tcaactaatg ttgcaaatct ttttggtgga 300 gaaggtgttc aagcggctcc agctcaagag ggtgcacaac aggagggagt tcaaccagct 360 ccagctcaag gtgggattag ctctccaatt aatgttacaa ctgctattga tgctaatgca 420 tcgcttacaa agattgaaga tgctattaga atggtaactg atcaaagagc aaatcttggt 480 gctttccaaa atagacttga gtctgttaaa gctagcacag attatgctat tgaaaactta 540 aaagcgtctt atgctcaaat taaagatgca ataatgacag atgaaattgt agcatctaca 600 accaacaqta ttttqacaca atctqcaatg gctatgattg c <210> 2 <211> 213 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide

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1 . 5 . 10 . 15

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Met His Met Leu Ser Asn Lys Ser Ser Ala Gln Asn Val Lys Thr Ala 35 40 45

Glu Glu Leu Gly Met Gln Pro Ala Lys Ile Asn Thr Pro Ala Ser Leu 50 55 60

Thr Gly Ala Gln Ala Ser Trp Thr Leu Arg Val Gln Val Gly Ala Asn
65 70 75 80

Gln Asp Glu Ala Ile Ala Val Asn Ile Phe Ser Thr Asn Val Ala Asn 85 90 95

Leu Phe Gly Gly Glu Gly Val Gln Ala Ala Pro Ala Gln Glu Gly Ala 100 105 110

Gln Gln Glu Gly Val Gln Pro Ala Pro Ala Gln Gly Gly Ile Ser Ser 115 120 125

Pro Ile Asn Val Thr Thr Ala Ile Asp Ala Asn Ala Ser Leu Thr Lys 130 135 140

Ile Glu Asp Ala Ile Arg Met Val Thr Asp Gln Arg Ala Asn Leu Gly
145 150 155 160

Ala Phe Gln Asn Arg Leu Glu Ser Val Lys Ala Ser Thr Asp Tyr Ala 165 170 175

Ile Glu Asn Leu Lys Ala Ser Tyr Ala Gln Ile Lys Asp Ala Ile Met 180 185 190

Thr Asp Glu Ile Val Ala Ser Thr Thr Asn Ser Ile Leu Thr Gln Ser 195 200 205

Ala Met Ala Met Ile 210

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gtaggaaatg acaaggtgat gacgttaatt tatgaataag ccccggctaa ttacgtgcca 480
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tccagctcaa gagggtgcac aacaggaggg agttcaacca gctccagctc aaggtgggat 240
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Ala Gln Ala Ala
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Ala Gln Thr Ala
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gctcaataca accagatgca tatgttatct aacaaatcat ctgctcaaaa tgtaaaaact 180
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aatattttct caactaatgt tgcaaatctt tttggtggag aaggtgttca agcggctcca 360
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Pro Ala Gln Gly Gly Ile Ser Ser Pro Ile Asn Val Thr Thr Ala Ile
Asp Ala Asn
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<211> 67
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Ile Phe Ser Thr Asn Val Ala Asn Leu Phe Gly Gly Glu Gly Val Gln
             20
                                 25
                                                     30
Ala Ala Pro Ala Gln Glu Gly Ala Gln Glu Gly Val Gln Pro Ala
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Pro Ala Gln Gly Gly Ile Ser Ser Pro Ile Asn Val Thr Thr Ala Ile
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Asp Ala Asn
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Ile Tyr Ala Ala Asn Val Ala Asn Leu Phe Ala Gly Glu Gly Ala Gln

0 25 30

Val Ser Pro Ala Gln Glu Gly Ala Gln Gln Glu Gly Val Gln Ala Ala 35 40 45

Pro Ala Pro Ala Ala Pro Ala Gln Gly Gly Val Asn Ser Pro Val 50 55 60

Asn Val Thr Thr Thr Ile Asp Ala Asn 65 70

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Ile Tyr Ala Ser Asn Val Ala Asn Leu Phe Ala Gly Glu Gly Ala Gln 20 25 30

Val Ser Pro Ala Gln Glu Gly Ala Gln Glu Gly Val Gln Ala Thr 35 40 45

Pro Ala Pro Val Ala Ala Pro Ala Pro Gly Gly Val Asn Ser Pro Ile 50 55 60

Asn Val Ile Thr Thr Val Asp Ala Asn 65 70

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Ile Tyr Ala Ala Asn Val Ala Asn Leu Phe Ala Gly Glu Gly Ala Gln 20 25 30

Ala Ala Pro Ala Gln Glu Gly Ala Gln Glu Gly Val Gln Ala Thr 35 40 45 Pro Ala Pro Val Ala Ala Pro Ala Pro Gly Gly Val Asn Ser Pro Ile 50 55 60

Asn Val Ile Thr Ala Val Asp Ala Asn 65 70

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<212> PRT

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Ile Tyr Ala Ser Asn Val Ala Asn Leu Phe Ala Gly Glu Gly Ala Gln
20 25 30

Ala Ala Pro Val Gln Glu Ile Gly Gln Gln Glu Glu Gly Gln Ala Ala 35 40 45

Pro Ala Pro Ala Ala Pro Ala Gln Gly Gly Val Asn Ser Pro Ile 50 55 60

Asn Val Thr Thr Ala Val Asp Ala Asn 65 70

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Ile Tyr Ala Ala Asn Val Ala Asn Leu Phe Ser Gly Glu Gly Ala Gln
20 25 30

Gln Val Ala Pro Ala Gln Glu Gly Ala Gln Gln Glu Gly Ala Gln Ala 35 40 45

Ala Pro Ala Pro Ala Ser Ala Pro Ala Gln Gly Gly Val Asn Ser Pro 50 55 60

Val Asn Val Thr Thr Ala Val Asp Ala Asn 65 70

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Ala Ala Gln Ala Ala Pro Val Gln Glu Gly Ala Gln Glu Gly Ala
35 40 45

Gln Gln Pro Thr Pro Ala Thr Ala Pro Thr Gln Gly Gly Val Asn Ser
50 55 60

Pro Val Asn Val Thr Thr Thr Val Asp Ala Asn 65 70 75

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Ile Tyr Ala Ala Asn Val Ala Asn Leu Phe Ser Gly Glu Gly Ala Gln 20 25 30

Ala Ala Gln Thr Ala Pro Val Gln Glu Gly Ala Gln Gln Glu Gly Ala 35 40 45

Gln Gln Pro Ala Pro Val Thr Ala Pro Ser Gln Gly Gly Val Asn Ser
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Pro Val Asn Val Thr Thr Thr Val Asp Ala Asn 65 70 75

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n Leu Phe Ser Gly Glu Gly Ala Gl
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Thr Ala Gln Ala Ala Pro Val Gln Glu Gly Val Gln Gln Glu Gly Ala 35 40 45

Gln Gln Pro Ala Pro Ala Thr Ala Pro Ser Gln Gly Gly Val Asn Ser 50 55 60

Pro Val Asn Val Thr Thr Thr Val Asp Ala Asn 65 70 75